## SEQUENCE LISTING

## (1) GENERAL INFORMATION

(i) APPLICANT: Koichi, Ishiguro Kazuki, Sato Jun-Mi, Park Tsuneko, Uchida Kazutomo Imahori

(ii) TITLE OF INVENTION: ANTI-PHOSPHORYLATED TAU PROTEIN ANTIBODY AND METHOD FOR DETECTING ALZHEIMER'S DISEASE WITH THE USE OF THE SAME

(iii) NUMBER OF SEQUENCES: 22

## (iv) COMPUTER READABLE FORM:

- (A) MEDIUM TYPE: Floppy disk
- (B) COMPUTER: IBM PC compatible
- (C) OPERATING SYSTEM: PC-DOS/MS-DOS
- (D) SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)
- (vi) CURRENT APPLICATION DATA
  - (A)APPLICATION NUMBER:
  - (B)FILING DATE:
  - (C)CLASSIFICATION:
- (vi) PRIOR APPLICATION DATA
  - (A)APPLICATION NUMBER: JP 8/56090
  - (B)FILING DATE: 13-MAR-1996
- (2) INFORMATION FOR SEQ ID NO: 1:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) SEQUENCE LENGTH: 441 amino acids
    - (B) SEQUENCE TYPE: amino acid
    - (D) TOPOLOGY:/linear
  - (ii) MOLECULE TYPE ≠ protein
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

Met Ala Glu Pro Arg Gin Glu Phe Glu Val Met Glu Asp His Ala Gly

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Gln Asp Thr Tyr Gly/Leu Gly Asp Arg Lys Asp Gln Gly Gly Tyr Thr

Met His Gln Glu Gly Asp Thr Asp Ala Gly Leu Lys Glu Ser Pro Leu

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35 40 45 Gln Thr Pro Thr Glu Asp Gly Ser Glu Glu Pro Gly Ser Glu Thr Ser 50 55 60 Asp Ala Lys Ser Thr Pro Thr Ala Glu Asp Val Thr Ala Pro Leu Val 75 Asp Glu Gly Ala Pro Gly Lys Gln Ala Ala Ala Gln Pro His Thr Glu 85 Ile Pro Glu Gly Thr Thr Ala Glu Glu Ala Gly Ile Gly Asp Thr Pro 105 Ser Leu Glu Asp Glu Ala Ala Gly His Val Thr Gln Ala Arg Met Val 115 120 Ser Lys Ser Lys Asp Gly Thr Gly Ser Asp Asp Lys Lys Ala Lys Gly 135 Ala Asp Gly Lys Thr Lys Ile Ala Thr Pro Arg Gly Ala Ala Pro Pro 150 160 Gly Gln Lys Gly Gln Ala Asn Ala Thr Arg Ile Pro Ala Lys Thr Pro 165 170 Pro Ala Pro Lys Thr Pro Pro Ser Ser Gly Glu Pro Pro Lys Ser Gly 185 Asp Arg Ser Gly Tyr Ser Ser Pro Gly Ser Pro Gly Thr Pro Gly Ser 200 Arg Ser Arg Thr Pro Ser Leu Pro Thr Pro Pro Thr Arg Glu Pro Lys 215 Lys Val Ala Val Val Arg Thr Pro Pro Lys Ser Pro Ser Ser Ala Lys 230 235 Ser Arg Leu Gln Thr Ala Pro Val Pro Met Pro Asp Leu Lys Asn Val 245 250 Lys Ser Lys Ile Gly Ser Thr Glu Asn Leu Lys His Gln Pro Gly Gly 260 265 Gly Lys Val Gln Ile Ile Asn Lys Leu Asp Leu Ser Asn Val Gln 275 280 285 Ser Lys Cys Gly Ser Lys Asp Asn Ile Lys His Val Pro Gly Gly Gly 295 Ser Val Gln Ile Val Tyr Lys Pro Val Asp Leu Ser Lys Val Thr Ser 305 310 315 320 Lys Cys Gly Ser Leu Gly Asn Ile His His Lys Pro Gly Gly Gln 330 Val Glu Val Lys Ser Glu Lys Leu Asp Phe Lys Asp Arg Val Gln Ser 340 345 350 Lys Ile Gly Ser Leu Asp Asn Ile Thr His Val Pro Gly Gly Asn

355 360 Lys Lys Ile Glu Thr His Lys Leu Thr Phe Arg Glu Asn Ala Lys Ala 370 375 380 Lys Thr Asp His Gly Ala Glu Ile Val Tyr Lys Ser Pro Val Val Ser 390 395 400 Gly Asp Thr Ser Pro Arg His Leu Ser Asn Val Ser Ser Thr Gly Ser 405 410 415 Ile Asp Met Val Asp Ser Pro Gln Leu Ala Thr Leu Ala Asp Glu Val 430 Ser Ala Ser Leu Ala Lys Gln Gly Leu 435 440

- (2) INFORMATION FOR SEQ ID NO: 2:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) SEQUENCE LENGTH: 12 amino acids
    - (B) SEQUENCE TYPE: amino acid
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (ix) FEATURE:
    - (A) NAME/KEY:
    - (B) LOCATION: 6
    - (D) OTHER INFORMATION: Xaa = phosphoserine
- (2) INFORMATION FOR SEQ ID NO: 3:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) SEQUENCE LENGTH: 13 amino acids
    - (B) SEQUENCE TYPE: amino acid
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (ix) FEATURE:
    - (A) NAME/KEY:
    - (B) LOCATION: 6
    - (D) OTHER INFORMATION: Xaa = phosphoserine

- (2) INFORMATION FOR SEQ ID NO: 4:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) SEQUENCE LENGTH: 12 amino acids
    - (B) SEQUENCE TYPE: amino acid
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (ix) FEATURE:
    - (A) NAME/KEY:
    - (B) LOCATION: 7
    - (D) OTHER INFORMATION: Xaa = phosphothreonine
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4: Cys Pro Gly Ser Pro Gly Xaa Pro Gly Ser Arg Ser

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- (2) INFORMATION FOR SEQ ID NO: 5:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) SEQUENCE LENGTH: 13 amino acids
    - (B) SEQUENCE TYPE: amino acid
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (ix) FEATURE:
    - (A) NAME/KEY:
    - (B) LOCATION: 6
    - (D) OTHER INFORMATION: Xaa = phosphoserine
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
  Lys Ser Xaa Pro Gly Xaa Pro Gly Thr Pro Gly Ser Arg
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- (2) INFORMATION FOR SEQ ID NO: 6:
  - (i) SEQUENCE CHARACTERISTICS:(A) SEQUENCE LENGTH: 14 amino acids
    - (B) SEQUENCE TYPE: amino acid
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (ix) FEATURE:
    - (A) NAME/KEY:
    - (B) LOCATION: 7
    - (D) OTHER INFORMATION: Xaa = phosphothreonine
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6: Cys Val Ala Val Val Arg Xaa Pro Pro Lys Ser Pro Ser Ser

- (2) INFORMATION FOR SEQ ID NO: 7:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) SEQUENCE LENGTH: 12 amino acids
    - (B) SEQUENCE TYPE: amino acid
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (ix) FEATURE:
    - (A) NAME/KEY:
    - (B) LOCATION: 7
    - (D) OTHER INFORMATION: Xaa = phosphoserine
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7: Cys Arg Thr Pro Pro Lys Xaa Pro Ser Ser Ala Lys 1 5 10
- (2) INFORMATION FOR SEQ ID NO: 8:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) SEQUENCE LENGTH: 12 amino acids
    - (B) SEQUENCE TYPE: amino acid
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (ix) FEATURE:
    - (A) NAME/KEY:
    - (B) LOCATION: 7
    - (D) OTHER INFORMATION: Xaa = phosphoserine
- (2) INFORMATION FOR SEQ ID NO: 9:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) SEQUENCE LENGTH: 12 amino acids
    - (B) SEQUENCE TYPE: amino acid
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (ix) FEATURE:
    - (A) NAME/KEY:
    - (B) LOCATION: 3
    - (D) OTHER INFORMATION: Xaa = phosphothreonine

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- (ix) FEATURE:
  - (A) NAME/KEY:
  - (B) LOCATION: 7
  - (D) OTHER INFORMATION: Xaa = phosphoserine
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:

Cys Arg Xaa Pro Pro Lys Xaa Pro Ser Ser Ala Lys

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- (2) INFORMATION FOR SEQ ID NO: 10:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) SEQUENCE LENGTH: 12 amino acids
    - (B) SEQUENCE TYPE: amino acid
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (ix) FEATURE:
    - (A) NAME/KEY:
    - (B) LOCATION: 7
    - (D) OTHER INFORMATION: Xaa = phosphoserine
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:

Cys Lys Ser Lys Ile Gly Xaa Thr Glu Asn Leu Lys

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(2) INFORMATION FOR SEQ ID NO: 11:

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- (i) SEQUENCE CHARACTERISTICS:
  - (A) SEQUENCE LENGTH: 12 amino acids
  - (B) SEQUENCE TYPE: amino acid
  - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
  - (A) NAME/KEY:
  - (B) LOCATION: 7
  - (D) OTHER INFORMATION: Xaa = phosphoserine
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:

Cys Glu Ile Val Tyr Lys Xaa Pro Val Val Ser Gly

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- (2) INFORMATION FOR SEQ ID NO: 12:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) SEQUENCE LENGTH: 12 amino acids
    - (B) SEQUENCE TYPE: amino acid

(D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
  - (A) NAME/KEY:
  - (B) LOCATION: 7
  - (D) OTHER INFORMATION: Xaa = phosphoserine
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12: Cys Val Ser Gly Asp Thr Xaa Pro Arg His Leu Ser

- (2) INFORMATION FOR SEQ ID NO: 13:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) SEQUENCE LENGTH: 12 amino acids
    - (B) SEQUENCE TYPE: amino acid
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (ix) FEATURE:
    - (A) NAME/KEY:
    - (B) LOCATION: 7
    - (D) OTHER INFORMATION: Xaa = phosphoserine
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13: Lys Leu Ser Asn Val Ser Xaa Thr Gly Ser Ile Asp 10
- (2) INFORMATION FOR SEQ ID NO: 14:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) SEQUENCE LENGTH: 12 amino acids
    - (B) SEQUENCE TYPE: amino acid
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (ix) FEATURE:
    - (A) NAME/KEY:
    - (B) LOCATION: 7
    - (D) OTHER INFORMATION: Xaa = phosphoserine
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14: Cys Ile Asp Met Val Asp Xaa Pro Gln Leu Ala Thr
- 5 10
- (2) INFORMATION FOR SEQ ID NO: 15:
  - (i) SEQUENCE CHARACTERISTICS:

- (A) SEQUENCE LENGTH: 12 amino acids(B) SEQUENCE TYPE: amino acid
- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
  - (A) NAME/KEY:
  - (B) LOCATION: 6
  - (D) OTHER INFORMATION:Xaa=phosphoserine
- (2) INFORMATION FOR SEQ ID NO: 16:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) SEQUENCE LENGTH: 12 amino acids
    - (B) SEQUENCE TYPE: amino acid
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (ix) FEATURE:
    - (A) NAME/KEY:
    - (B) LOCATION: 6
    - (D) OTHER INFORMATION: Xaa = phosphoserine
  - (ix) FEATURE:
    - (A) NAME/KEY:
    - (B) LOCATION: 7
    - (D) OTHER INFORMATION: Xaa = phosphoserine
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16: Lys Leu Ser Asn Val Xaa Xaa Thr Gly Ser Ile Asp

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- (2) INFORMATION FOR SEQ ID NO: 17:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) SEQUENCE LENGTH: 17 amino acids
    - (B) SEQUENCE TYPE: amino acid
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:

Ser Pro Gln Leu Ala Thr Leu Ala Asp Glu Val Ser Ala Ser Leu Ala Lys

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(2)	INFORMATION FOR SEQ ID NO: 18:  (i) SEQUENCE CHARACTERISTICS:
	<ul><li>(A) SEQUENCE LENGTH: 12 amino acids</li><li>(B) SEQUENCE TYPE: amino acid</li><li>(D) TOPOLOGY: linear</li></ul>
	(ii) MOLECULE TYPE: peptide
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:
Ala	Glu Pro Arg Gln Glu Phe Glu Val Met Glu Cys
1	5 10
(2)	INFORMATION FOR SEQ ID NO: 19: (i) SEQUENCE CHARACTERISTICS:
	(A) SEQUENCE LENGTH: 15 amino acids
	(B) SEQUENCE TYPE: amino acid
	(D) TOPOLOGY: linear
	(ii) MOLECULE TYPE: peptide
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19:
Val	Ala Val Val Arg Thr Pro Pro Lys Ser Pro Ser Ser Ala Lys
1	5 10 15
(2)	INFORMATION FOR SEQ ID NO: 20:
	(i) SEQUENCE CHARACTERISTICS:
	(A) SEQUENCE LENGTH: 34 amino acids
	(B) SEQUENCE TYPE: amino acid
	(D) TOPOLOGY: linear
	(ii) MOLECULE TYPE: peptide
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:
	Gly Asp Arg Ser Gly Tyr Ser Ser Pro Gly Ser Pro Gly Thr Pro
1	5 10 15
GIY	Ser Arg Ser Arg Thr Pro Ser Leu Pro Thr Pro Pro Thr Arg Glu
D	20 25 30
Pro	ràs
(2)	INFORMATION FOR SEQ ID NO: 21:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) SEQUENCE LENGTH: 55 amino acids
  - (B) SEQUENCE TYPE: amino acid
  - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 21:

Ala Lys Thr Asp His Gly Ala Glu Ile Val Tyr Lys Ser Pro Val Val

1 5 10 15

Ser Gly Asp Thr Ser Pro Arg His Leu Ser Asn Val Ser Ser Thr Gly 20 25 30

Ser Ile Asp Met Val Asp Ser Pro Gln Leu Ala Thr Leu Ala Asp Glu 35 40 45

Val Ser Ala Ser Leu Ala Lys 50 55

- (2) INFORMATION FOR SEQ ID NO: 22:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) SEQUENCE LENGTH: 12 amino acids
    - (B) SEQUENCE TYPE: amino acid
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide

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